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0814

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/915814

CRF Processing Date: 10/09/01
Edited by: MH
Verified by: _____ (ST

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: _____

☐ Edited the Current Application Data section with the actual current number. The number inputted by tH applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integ

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: ENTERED

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: _____

☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end c ☐ page numbers throughout text; ☐ other invalid text, such as _____

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically: _____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (er due to a PatentIn bug). Sequences corrected: _____

☐ Other: _____

_____

_____

*Examiner: The above corrections must be communicated to the applicant in the first Offic Action. DO NOT send a copy of this form.

3/1/9

```
          RAW SEQUENCE LISTING              DATE: 10/09/2001
          PATENT APPLICATION:  US/09/915,814     TIME: 10:40:24

          Input Set : A:\PTO.MH.txt
          Output Set: N:\CRF3\10092001\I915814.raw

   3 <110> APPLICANT: Madeline M. Butler
   4        Andrew T. Watt
   5        Susan M. Freier
   6        Jacqueline Wyatt
   8 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF HORMONE-SENSITIVE LIPASE EXPRESSION
  10 <130> FILE REFERENCE: ISPH-0587
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/915,814
C--> 12 <141> CURRENT FILING DATE: 2001-07-26
  12 <160> NUMBER OF SEQ ID NOS: 230
  14 <210> SEQ ID NO: 1
  15 <211> LENGTH: 20
  16 <212> TYPE: DNA
  17 <213> ORGANISM: Artificial Sequence
  19 <220> FEATURE:
  20 <223> OTHER INFORMATION: Antisense Oligonucleotide
  22 <400> SEQUENCE: 1
  23 tccgtcatcg ctcctcaggg                                        20
  25 <210> SEQ ID NO: 2
  26 <211> LENGTH: 20
  27 <212> TYPE: DNA
  28 <213> ORGANISM: Artificial Sequence
  30 <220> FEATURE:
  31 <223> OTHER INFORMATION: Antisense Oligonucleotide
  33 <400> SEQUENCE: 2
  34 atgcattctg cccccaagga                                        20
  36 <210> SEQ ID NO: 3
  37 <211> LENGTH: 3804
  38 <212> TYPE: DNA
  39 <213> ORGANISM: Homo sapiens
  41 <220> FEATURE:
  42 <221> NAME/KEY: CDS
  43 <222> LOCATION: (278)...(3508)
  45 <400> SEQUENCE: 3
  46 cttcttgtaa gagagtgcta ggcacatagc cccctcctat tcctaatcct cccaccaaag 60
  47 aaagaggcac agagttcatt acttagtggg ggccagctgt gatcggccaa ctgccagctg 120
  48 ccttaaaaag gaagaccagt gatgctagga tggagtgaaa cccaagagga agtgccatca 180
  49 tgaggaatca atgagagatc tgtgaagaga gagggctggg tgggagccca gaaggataga 240
  50 acctggaaga tcaatatctc ccgtgaggga aataaca atg gag cca ggt tct aag  295
  51                                             Met Glu Pro Gly Ser Lys
  52                                              1               5
  54 tca gtg tct agg tca gac tgg caa cct gaa cca cac cag agg cct ata   343
  55 Ser Val Ser Arg Ser Asp Trp Gln Pro Glu Pro His Gln Arg Pro Ile
  56              10              15                  20
  58 acc ccg cta gag cct ggg cca gaa aag aca ccc ata gcc cag cca gaa   391
  59 Thr Pro Leu Glu Pro Gly Pro Glu Lys Thr Pro Ile Ala Gln Pro Glu
  60          25              30                  35
  62 tcg aag act ctg cag gga tcc aat acc caa cag aag cct gct tca aac   439
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ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/915,814

DATE: 10/09/2001
TIME: 10:40:24

Input Set : A:\PTO.MH.txt
Output Set: N:\CRF3\10092001\I915814.raw

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63 Ser Lys Thr Leu Gln Gly Ser Asn Thr Gln Gln Lys Pro Ala Ser Asn
64      40              45              50
66 caa aga ccc ctc acc cag cag gag acc cct gca caa cat gat gct gaa      487
67 Gln Arg Pro Leu Thr Gln Gln Glu Thr Pro Ala Gln His Asp Ala Glu
68 55              60              65              70
70 tcc cag aag gaa cct aga gcc caa caa aaa tct gct tca caa gag gaa      535
71 Ser Gln Lys Glu Pro Arg Ala Gln Gln Lys Ser Ala Ser Gln Glu Glu
72              75              80              85
74 ttt ctt gcc cca cag aag ccc gca cca cag caa tca cct tac atc caa      583
75 Phe Leu Ala Pro Gln Lys Pro Ala Pro Gln Gln Ser Pro Tyr Ile Gln
76              90              95              100
78 agg gtg ctg ctc act caa cag gaa gct gcc tcc cag cag gga cct ggg      631
79 Arg Val Leu Leu Thr Gln Gln Glu Ala Ala Ser Gln Gln Gly Pro Gly
80              105              110              115
82 cta gga aaa gaa tct ata act caa cag gag cca gca ttg aga caa aga      679
83 Leu Gly Lys Glu Ser Ile Thr Gln Gln Glu Pro Ala Leu Arg Gln Arg
84      120              125              130
86 cat gta gcc cag cca ggg cct ggg cca gga gag cca cct cca gct caa      727
87 His Val Ala Gln Pro Gly Pro Gly Pro Gly Glu Pro Pro Pro Ala Gln
88 135              140              145              150
90 caa gaa gct gaa tca aca cct gcg gcc cag gct aaa cct gga gcc aaa      775
91 Gln Glu Ala Glu Ser Thr Pro Ala Ala Gln Ala Lys Pro Gly Ala Lys
92              155              160              165
94 agg gag cca tct gcc ccg act gaa tct aca tcc caa gag aca cct gaa      823
95 Arg Glu Pro Ser Ala Pro Thr Glu Ser Thr Ser Gln Glu Thr Pro Glu
96              170              175              180
98 cag tca gac aag caa aca acg cca gtc cag gga gcc aaa tcc aag cag      871
99 Gln Ser Asp Lys Gln Thr Thr Pro Val Gln Gly Ala Lys Ser Lys Gln
100              185              190              195
102 gga tct ttg aca gag ctg gga ttt cta aca aaa ctt cag gaa cta tcc      919
103 Gly Ser Leu Thr Glu Leu Gly Phe Leu Thr Lys Leu Gln Glu Leu Ser
104      200              205              210
106 ata cag cga tca gcc cta gag tgg aag gca ctt tct gag tgg gtc gca      967
107 Ile Gln Arg Ser Ala Leu Glu Trp Lys Ala Leu Ser Glu Trp Val Ala
108 215              220              225              230
110 gat tct gag tca gaa tca gat gtg gga tca tct tca gac aca gat tct      1015
111 Asp Ser Glu Ser Glu Ser Asp Val Gly Ser Ser Ser Asp Thr Asp Ser
112              235              240              245
114 cca gcc acg atg ggt gga atg gtg gcc cag gga gtg aag cta ggc ttc      1063
115 Pro Ala Thr Met Gly Gly Met Val Ala Gln Gly Val Lys Leu Gly Phe
116              250              255              260
118 aaa gga aaa tct ggt tat aaa gtg atg tca gga tac agt ggg acg tcg      1111
119 Lys Gly Lys Ser Gly Tyr Lys Val Met Ser Gly Tyr Ser Gly Thr Ser
120              265              270              275
122 cca cat gag aaa acc agt gct cgg aat cac aga cac tac cag gat aca      1159
123 Pro His Glu Lys Thr Ser Ala Arg Asn His Arg His Tyr Gln Asp Thr
124      280              285              290
126 gcc tca agg ctc atc cac aac atg gac ctg cgc aca atg aca cag tcg      1207
127 Ala Ser Arg Leu Ile His Asn Met Asp Leu Arg Thr Met Thr Gln Ser
```

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128 295                     300                    305                      310
130 ctg gtg act ctg gcg gag gac aac ata gcc ttc ttc tcg agc cag ggt    1255
131 Leu Val Thr Leu Ala Glu Asp Asn Ile Ala Phe Phe Ser Ser Gln Gly
132                 315                     320                     325       .
134 cct ggg gaa acg gcc cag cgg ctg tca ggc gtt ttt gcc ggt gta cgg    1303
135 Pro Gly Glu Thr Ala Gln Arg Leu Ser Gly Val Phe Ala Gly Val Arg     .
136                 330                     335                     340
138 gag cag gcg ctg ggg ctg gag ccg gcc ctg ggc cgc ctg ctg ggt gtg    1351
139 Glu Gln Ala Leu Gly Leu Glu Pro Ala Leu Gly Arg Leu Leu Gly Val
140         345              .      350                     355
142 gcg cac ctc ttt gac ctg gac cca gag aca ccg gcc aac ggg tac cgc    1399
143 Ala His Leu Phe Asp Leu Asp Pro Glu Thr Pro Ala Asn Gly Tyr Arg
144     360                     365                     370
146 agc cta gtg cac aca gcc cgc tgc tgc ctg gcg cac ctc ctg cac aaa    1447
147 Ser Leu Val His Thr Ala Arg Cys Cys Leu Ala His Leu Leu His Lys
148 375                     380                     385                      390
150 tcc cgc tat gtg gcc tcc aac cgc cgc agc atc ttc ttc cgc acc agc    1495
151 Ser Arg Tyr Val Ala Ser Asn Arg Arg Ser Ile Phe Phe Arg Thr Ser
152                 395                     400                     405
154 cac aac ctg gcc gag ctg gag gcc tac ctg gct gcc ctc acc cag ctc    1543
155 His Asn Leu Ala Glu Leu Glu Ala Tyr Leu Ala Ala Leu Thr Gln Leu
156                 410                     415                     420
158 cgc gct ctg gtc tac tac gcc cag cgc ctg ctg gtt acc aat cgg ccg    1591
159 Arg Ala Leu Val Tyr Tyr Ala Gln Arg Leu Leu Val Thr Asn Arg Pro
160         425                     430                     435
162 ggg gta ctc ttc ttt gag ggc gac gag ggg ctc acc gcc gac ttc ctc    1639
163 Gly Val Leu Phe Phe Glu Gly Asp Glu Gly Leu Thr Ala Asp Phe Leu
164     440                     445                     450
166 cgg gag tat gtc acg ctg cat aag gga tgc ttc tat ggc cgc tgc ctg    1687
167 Arg Glu Tyr Val Thr Leu His Lys Gly Cys Phe Tyr Gly Arg Cys Leu
168 455                     460                     465                      470
170 ggc ttc cag ttc acg cct gcc atc cgg cca ttc ctg cag acc atc tcc    1735
171 Gly Phe Gln Phe Thr Pro Ala Ile Arg Pro Phe Leu Gln Thr Ile Ser
172                 475                     480                     485
174 att ggg ctg gtg tcc ttc ggg gag cac tac aaa cgc aac gag aca ggc    1783
175 Ile Gly Leu Val Ser Phe Gly Glu His Tyr Lys Arg Asn Glu Thr Gly
176                 490                     495                     500
178 ctc agt gtg gcc gcc agc tct ctc ttc acc agc ggc cgc ttt gcc atc    1831
179 Leu Ser Val Ala Ala Ser Ser Leu Phe Thr Ser Gly Arg Phe Ala Ile
180         505                     510                     515
182 gac ccc gag ctg cgt ggg gct gag ttt gag cgg atc aca cag aac ctg    1879
183 Asp Pro Glu Leu Arg Gly Ala Glu Phe Glu Arg Ile Thr Gln Asn Leu
184     520                     525                     530
186 gac gtg cac ttc tgg aaa gcc ttc tgg aac atc acc gag atg gaa gtg    1927
187 Asp Val His Phe Trp Lys Ala Phe Trp Asn Ile Thr Glu Met Glu Val
188 535                     540                     545                      550
190 cta tcg tct ctg gcc aac atg gca tcg gcc acc gtg agg gta agc cgc    1975
191 Leu Ser Ser Leu Ala Asn Met Ala Ser Ala Thr Val Arg Val Ser Arg
192                 555                     560                     565
```

```
194 ctg ctc agc ctg cca ccc gaa gcc ttt gag atg cca ctg act gcc gac    2023
195 Leu Leu Ser Leu Pro Pro Glu Ala Phe Glu Met Pro Leu Thr Ala Asp
196         570             575             580
198 ccc acg ctc acg gtc acc atc tca ccc cca ctg gcc cac aca ggc cct    2071
199 Pro Thr Leu Thr Val Thr Ile Ser Pro Pro Leu Ala His Thr Gly Pro
200         585             590             595
202 ggg ccc gtc ctc gtc agg ctc atc tcc tat gac ctg cgt gaa gga cag    2119
203 Gly Pro Val Leu Val Arg Leu Ile Ser Tyr Asp Leu Arg Glu Gly Gln
204     600             605                 610
206 gac agt gag gag ctc agc agc ctg ata aag tcc aac ggc caa cgg agc    2167
207 Asp Ser Glu Glu Leu Ser Ser Leu Ile Lys Ser Asn Gly Gln Arg Ser
208 615             620                 625                 630
210 ctg gag ctg tgg ccg cgc ccc cag cag gca ccc cgc tcg cgg tcc ctg    2215
211 Leu Glu Leu Trp Pro Arg Pro Gln Gln Ala Pro Arg Ser Arg Ser Leu
212             635             640                 645
214 ata gtg cac ttc cac ggc ggt ggc ttt gtg gcc cag acc tcc aga tcc    2263
215 Ile Val His Phe His Gly Gly Gly Phe Val Ala Gln Thr Ser Arg Ser
216         650             655             .   660
218 cac gag ccc tac ctc aag agc tgg gcc cag gag ctg ggc gcc ccc atc    2311
219 His Glu Pro Tyr Leu Lys Ser Trp Ala Gln Glu Leu Gly Ala Pro Ile
220         665             670             675
222 atc tcc atc gac tac tcc ctg gcc cct gag gcc ccc ttc ccc cgt gcg    2359
223 Ile Ser Ile Asp Tyr Ser Leu Ala Pro Glu Ala Pro Phe Pro Arg Ala
224     680             685                 690
226 ctg gag gag tgc ttc ttc gcc tac tgc tgg gcc atc aag cac tgc gcc    2407
227 Leu Glu Glu Cys Phe Phe Ala Tyr Cys Trp Ala Ile Lys His Cys Ala
228 695             700             705                 710
230 ctc ctt ggc tca aca ggg gaa cga atc tgc ctt gcg ggg gac agt gca    2455
231 Leu Leu Gly Ser Thr Gly Glu Arg Ile Cys Leu Ala Gly Asp Ser Ala
232             715             720                 725
234 ggc ggg aac ctc tgc ttc acc gtg gct ctt cgg gca gca gcc tac ggg    2503
235 Gly Gly Asn Leu Cys Phe Thr Val Ala Leu Arg Ala Ala Ala Tyr Gly
236         730             735             740
238 gtg cgg gtg cca gat ggc atc atg gca gcc tac ccg gcc aca atg ctg    2551
239 Val Arg Val Pro Asp Gly Ile Met Ala Ala Tyr Pro Ala Thr Met Leu
240     ..  745             750             755
242 cag cct gcc gcc tct ccc tcc cgc ctg ctg agc ctc atg gac ccc ttg    2599
243 Gln Pro Ala Ala Ser Pro Ser Arg Leu Leu Ser Leu Met Asp Pro Leu
244     760             765             770 .
246 ctg ccc ctc agt gtg ctc tcc aag tgt gtc agc gcc tat gct ggt gca    2647
247 Leu Pro Leu Ser Val Leu Ser Lys Cys Val Ser Ala Tyr Ala Gly Ala
248 775             780             785                 790
250 aag acg gag gac cac tcc aac tca gac cag aaa gcc ctc ggc atg atg    2695
251 Lys Thr Glu Asp His Ser Asn Ser Asp Gln Lys Ala Leu Gly Met Met
252             795             800             805
254 ggg ctg gtg cgg cgg gac aca gcc ctg ctc ctc cga gac ttc cgc ctg    2743
255 Gly Leu Val Arg Arg Asp Thr Ala Leu Leu Leu Arg Asp Phe Arg Leu
256             810             815                 820
258 ggt gcc tcc tca tgg ctc aac tcc ttc ctg gag tta agt ggg cgc aag    2791
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RAW SEQUENCE LISTING   DATE: 10/09/2001
PATENT APPLICATION: US/09/915,814   TIME: 10:40:24

Input Set : A:\PTO.MH.txt
Output Set: N:\CRF3\10092001\I915814.raw

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259 Gly Ala Ser Ser Trp Leu Asn Ser Phe Leu Glu Leu Ser Gly Arg Lys
260         825             830             835
262 tcc cag aag atg tcg gag ccc ata gca gag ccg atg cgc cgc agt gtg   2839
263 Ser Gln Lys Met Ser Glu Pro Ile Ala Glu Pro Met Arg Arg Ser Val
264     840             845             850
266 tct gaa gca gca ctg gcc cag ccc cag ggc cca ctg ggc acg gat tcc   2887
267 Ser Glu Ala Ala Leu Ala Gln Pro Gln Gly Pro Leu Gly Thr Asp Ser
268 855             860             865                     870
270 ctc aag aac ctg acc ctg agg gac ttg agc ctg agg gga aac tcc gag   2935
271 Leu Lys Asn Leu Thr Leu Arg Asp Leu Ser Leu Arg Gly Asn Ser Glu
272             875             880             885
274 acg tcg tcg gac acc ccc gag atg tcg ctg tca gct gag aca ctt agc   2983
275 Thr Ser Ser Asp Thr Pro Glu Met Ser Leu Ser Ala Glu Thr Leu Ser
276             890             895             900
278 ccc tcc aca ccc tcc gat gtc aac ttc tta tta cca cct gag gat gca   3031
279 Pro Ser Thr Pro Ser Asp Val Asn Phe Leu Leu Pro Pro Glu Asp Ala
280         905             910             915
282 ggg gaa gag gct gag gcc aaa aat gag ctg agc ccc atg gac aga ggc   3079
283 Gly Glu Glu Ala Glu Ala Lys Asn Glu Leu Ser Pro Met Asp Arg Gly
284     920             925             930
286 ctg ggc gtc cgt gcc gcc ttc ccc gag ggt ttc cac ccc cga cgc tcc   3127
287 Leu Gly Val Arg Ala Ala Phe Pro Glu Gly Phe His Pro Arg Arg Ser
288 935             940             945                     950
290 agc cag ggt gcc aca cag atg ccc ctc tac tcc tca ccc ata gtc aag   3175
291 Ser Gln Gly Ala Thr Gln Met Pro Leu Tyr Ser Ser Pro Ile Val Lys
292             955             960             965
294 aac ccc ttc atg tcg ccg ctg ctg gca ccc gac agc atg ctc aag agc   3223
295 Asn Pro Phe Met Ser Pro Leu Leu Ala Pro Asp Ser Met Leu Lys Ser
296             970             975             980
298 ctg cca cct gtg cac atc gtg gcg tgc gcg ctg gac ccc atg ctg gac   3271
299 Leu Pro Pro Val His Ile Val Ala Cys Ala Leu Asp Pro Met Leu Asp
300         985             990             995
302 gac tcg gtc atg ctc gcg cgg cga ctg cgc aac ctg ggc cag ccg gtg   3319
303 Asp Ser Val Met Leu Ala Arg Arg Leu Arg Asn Leu Gly Gln Pro Val
304     1000            1005            1010
306 acg ctg cgc gtg gtg gag gac ctg ccg cac ggc ttc ctg acc cta gcg   3367
307 Thr Leu Arg Val Val Glu Asp Leu Pro His Gly Phe Leu Thr Leu Ala
308 1015            1020            1025                    1030
310 gcg ctg tgc cgc gag acg cgc cag gcc gca gag ctg tgc gtg gag cgc   3415
311 Ala Leu Cys Arg Glu Thr Arg Gln Ala Ala Glu Leu Cys Val Glu Arg
312             1035            1040            1045
314 atc cgc ctc gtc ctc act cct ccc gcc gga gcc ggg ccg agc ggg gag   3463
315 Ile Arg Leu Val Leu Thr Pro Pro Ala Gly Ala Gly Pro Ser Gly Glu
316             1050            1055            1060
318 acg ggg gct gcg ggg gta gac ggg ggc tgc ggg ggg cga cac taa       3508
319 Thr Gly Ala Ala Gly Val Asp Gly Gly Cys Gly Gly Arg His
320             1065            1070            1075
322 aagcctgttg ttcccatctg cgccggcctc cgtcatgaat gccttccggg ccgggcggaa 3568
323 ggggacgcgg gctgtgctta cttaagtcgg gggtggcaag ggggcggggc ggggccgaa  3628
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VERIFICATION SUMMARY   DATE: 10/09/2001
PATENT APPLICATION: **US/09/915,814**   TIME: 10:40:25

Input Set : **A:\PTO.MH.txt**
Output Set: **N:\CRF3\10092001\I915814.raw**

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date